

09824870-061304

>33358 cDNA ORF 75-1046  
GCGTCCGCGGACGCGTGGGTTATAACTCAGTGAAATTTTACAGTCCTAGGACCCTATACAGAGCATAAGC  
CAAAATGGAAGATGGTCCTGTTTTCTATGGCTTTAAAAACATTTTTATTACAATGTTTGCTACGTTTTTTT  
TTCTTTAAGCTTTTAATTAAAGTTTTTTTTGGCTCTCCTAACCATTCTATATCGTCAAAGGAAATAGAA  
AAGAAGCGGCTAGGATAGCAGAAGAGATCTATGGTGGAATTTTCAAGATTGCTGGGCTGATCGATCCCCACT  
TCATGAAGCTGCAGCTCAGGGGCGCTTACTGGCCCTTAAACTTTAATTGCACAAGGTGTCAATGTGAAC  
CTTGTGACAATTAACCGGGTGTCTTCTCTCCACGAGGCATGCCTTGGAGGTACAGTGGCCTGTGCCAAAG  
CCTTATTGGAAAATGGTGCACACGTCAATGGAGTGACAGTTTACGGAGCCACACCCCTCTTCAATGCTTG  
CTGCAGCGGCAGTGTGTCATGTGTCAATGTGCTGCTGGAGTTCGGAGCCAAGGCCAGTTGGAGGTGCAC  
CTGGCCTCGCCCATCCATGAGGCAGTGAAGAGAGGTACAGAGAGTGCATGGAGATCCTGCTGGCAAATA  
ATGTTAACATTGACCATGAGGTGCCCTCAGCTCGGAACCTCCCTATATGTGGCCTGCACCTACCAGAGGGT  
AGACTGTGTGAAGAACTTCTAGAATTAGGAGCCAGTGTGACCATGGCCAGTGGCTGGACACCCCACTC  
CATGCTGCAGCGAGGCAGTCCAATGTGGAGGTATCCACCTGCTAACCGACTATGGAGCTAACCTGAAGC  
GTAGAAATGCTCAGGGCAAAAGTGCCTTGATCTGGCGGCTCCAAAAGCAGCGTGGAGCAGGCACCTCTT  
GCTCCGTGAAGGCCACCTGCTCTTTCCAGCTCTGCCGCTGTGTGTCCGGAAGTGTCTCGGTGAGCA  
TGTCATCAAGCCATCCACAAGCTACATCTGCCAGAGCCACTCGA  
ACGATTCCTCCTATACCAATAGTCCTAAGTGTTCCTGGGAAGATACTTGAATGACACAGATTGTTGTCT  
GCTGTACCTAGAGTACCTAATGTAGAAGCTCAACAGCTTAGACTCCTAGTATCTTTAAATGAGMTCAGTC  
GAAGTAAATCCCCCATGAGCTAGAACACTTGAGGAGTGGRAACTCCTGGTTAGTTTAAATGTTCTCATTA  
CCAAGGGGCAAGTAGAAACCATTTAGCTTTTAGCTCTTTGTTGTTAAGAACTTAAAGAAGTGTGAAGT  
AGAGTGAAGAACAAATAGGCTGTTTTTTGATGATTTCGGGATCTTCTTGTACCTAAAAGTCAACATTCTGAAT  
ATTGTATAGACACATATAAATTCAGGTGGATAAGATTATAACAAATGTTAGGTATTCCAAGATATGttct  
tgatttagttccttccttcagcccttccccactttttttctttcttTCCTTGAATAAATCTGGTATAATT  
TTGAAAAAAAAAAAAAAAAAAAAA

Figure 1A

>33358 amino acids

MEDGPVFYGFKNIFITMFATFFFFKLLIKVFLALLTHFYIVKGNRKEAARIAEEIYGGISDCWADRSPLH  
EAAQGRLLALKTLIAQGVNVLVTINRVSSLHEACLGGHVACAKALLENGAHVNGVTVHGATPLFNACC  
SGSAACVNVLLFEGAQLEVLASPIHEAVKRGHRECMEILLANNVNIDHEVPQLGTPLYVACTYQRVD  
CVKKLLELGASVDHGQWLDTPLHAAARQSNVEVIHLLTDYGANLKRRNAQGKSALDLAAPKSSVEQALLL  
REGPPALSQLCRLCVRKCLGRACHQAIHKLHLPEPLERFLLYQ.

**Figure 1B**

09884870-061804

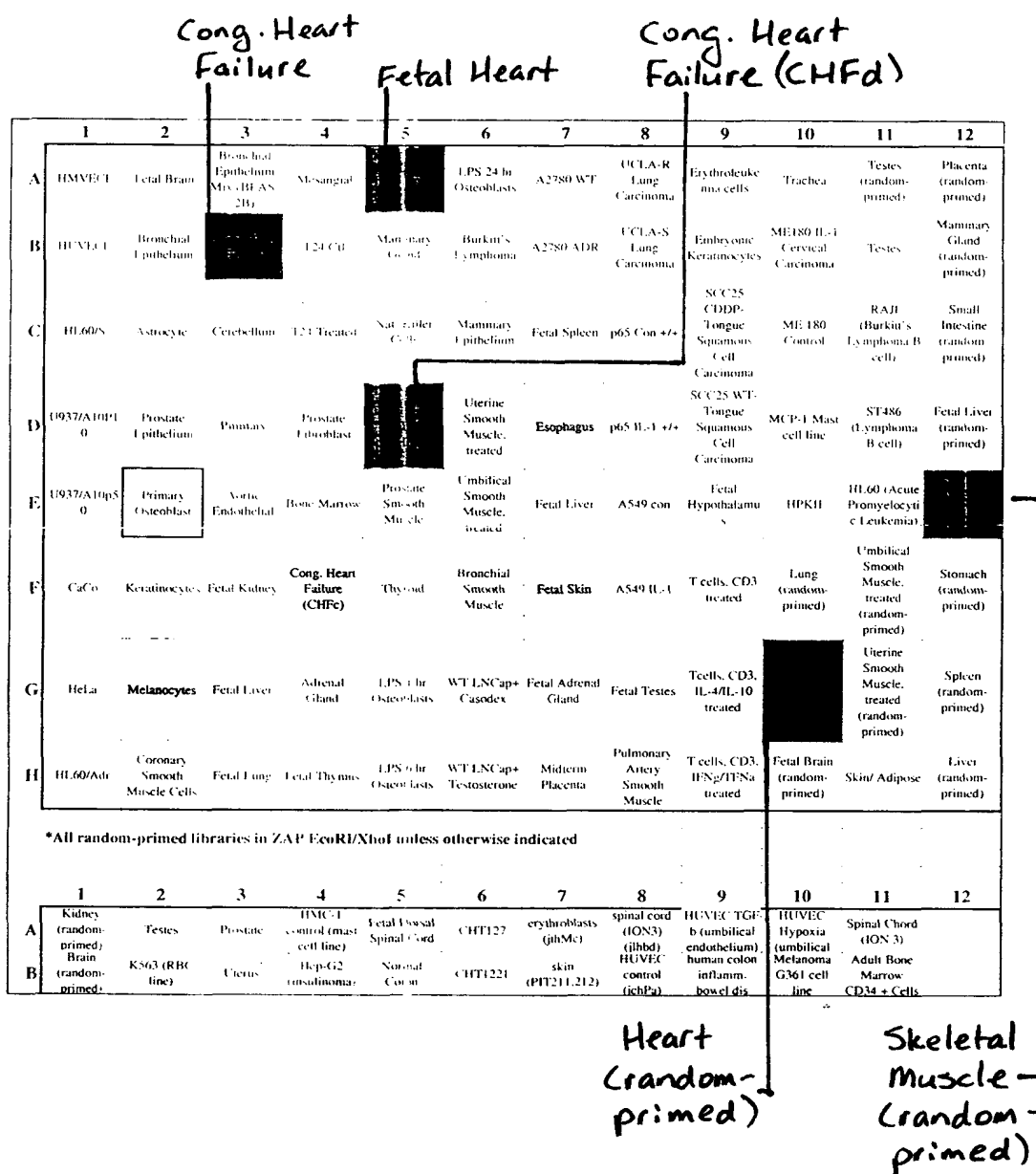


Figure 2A

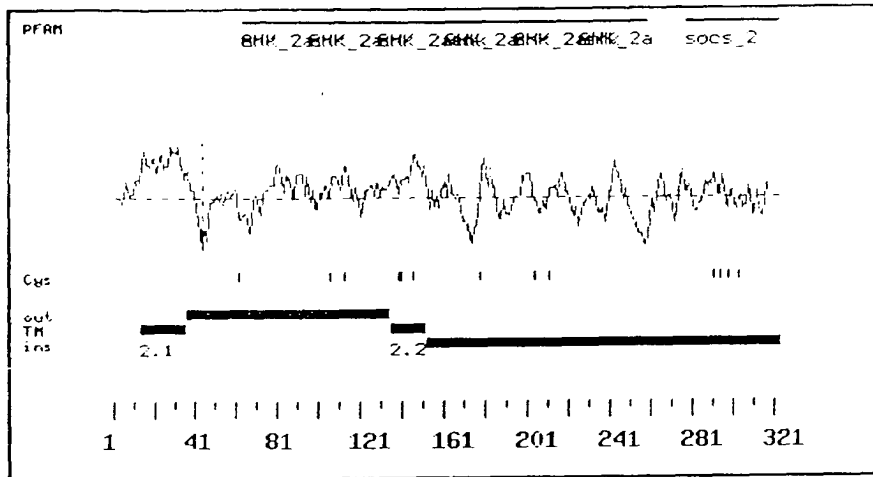
**Heart**

<b>C</b>	HPK (random-primed)	Lung	Mammary Gland	Normal Breast Epithelia	Colon to Liver Metastasis (CHT128)	normal breast	BMC1234 (jhbMc)	W138 20h serum starve emb. Lung	Th1 Cells	HUVEC untreated (umbilical endothelia)
<b>D</b>	Uterus (random-primed)	Liver	Spleen	Normal Ovarian Epithelia	Colon to Liver Metastasis (CHT133)	PTH Osteo	ovarian ascites (johOb)	lung squamous cell carcinoma	Th2 Cells	IBD Colon (WUM 23)
<b>E</b>	Thymus (random-primed)	Heart	Small Intestine	Normal Megakaryocytes	Colon Carcinoma (NDR109)	lung adenocarcinoma (PTT245)	IBD colon (WUM6) (jblina)	brain subcortical white matter	Prostate Tumor Xenograft A12	Trigeminal Ganglia
<b>F</b>	9 week Fetus	Thymus	Retinal Pigmentosa Epithelia	Bone Marrow	Colon Carcinoma (NDR103)	lung squamous cell carcinoma PTT299	Cervical Cancer (johCe)	normal prostate (ziploxs)	Prostate Tumor Xenograft K10	Lumbosacral Spinal Chord
<b>G</b>	A549 control (random-primed)	Stomach	Retina	Th1 induced T cell	Colon Carcinoma (NDRR2)	d8 dendritic cells	Spinal cord (jlbhc)	ovarian epithelium tumor	Prostate Cancer to Liver Metastasis JH13	Lumbosacral Dorsal Root Ganglia
<b>H</b>	Salivary Gland	Skeletal Muscle	HMC-1 (mast cell line)	Th2 induced T cell	Colon Carcinoma (NDR097)	Megakaryocytes (jhbcb)	DRG (ION6,7,8) (jhbce)	HUVEC name (umbilical end.)	Prostate Cancer to Liver Metastasis JH14	Dorsal Root Ganglia (ION 6,7 & 8)

**Figure 2B**

09884870 061801

## Analysis of 33358 (323 aa)



>33358  
MEDGPVFGFKNIFITMFATFFFFKLLIKVFLALLTHFYIVKGNRKEAARIAEEIYGGIS  
DCWADRSPLHEAAAQGRLLALKTLIAQGVNVNLVTINRVSSLHEACLGGHVACAKALLEN  
GAHVNGVTVHGATPLFNACCSGSAACVNVILLEFGAKAQLVHLASPIHEAVKRGHRECME  
ILLANNVNIDHEVPQLGTPLYVACTYQRVDCVKKLLELGASVDHGQWLDTPLHAAARQSN  
VEVIHLLTDYGANLKRRNAQGKSALDLAAPKSSVEQALLLREGPPALSQLCRLCVRKCLG  
RACHQAIHKLHLPEPLERFLLYQ

FIGURE 3